

Fig. 1

SEQ ID NO: 1

TCCTGGTCATCGATGTCATCCACGAGGTGGCCCACAGTTGGTTCGGCAACGC
TGTCACCAACGCCACGTGGGAAGAGATGTGGCTGAGCGAGGGCCTGGCCACC
TATGCCCAGCGCCGTATCACCACCGAAACCTACGGTGCTGCCTTCACCTGCC
TGGAGACTGCCTTCCGCCTGGACGCCCTGCACCGGCAGATGAAGCTTCTGGG
AGAGGACAGCCCGGTCAGCAAACCTGCAGGTCAAGCTGGAGCCAGGAGTGAAT
CCCAGCCACCTGATGAACCTGTTACCTACGAGAAGGGCTACTGCTTCGTGT
ACTACCTGTCCCAGCTCTGCGGAGACCCACAGCGCTTTGATGACTTTCTCCG
AGCCTATGTGGAGAAGTACAAGTTCACCAGCGTGGTGGCCCAGGACCTGCTG
GACTCCTTCCTGAGCTTCTTCCCGGAGCTGAAGGAGCAGAGCGTGGACTGCC
GGGCAGGGCTGGAATTCGAGCGCTGGCTCAATGCCACAGGCCCGCCGCTGGC
TGAGCCGGACCTGTCTCAGGGATCCAGCCTGACCCGGCCCGTGGAGGCCCTT
TTCCAGCTGTGGACCGCAGAACCTCTGGACCAGGCAGCTGCCTCGGCCAGCG
CCATTGACATCTCCAAGTGGAGGACCTTCAGACAGCACTCTTCCTGGACCG
GCTCCTGGATGGGTCCCCGCTGCCGCAGGAGGTGGTGATGAGCCTGTCCAAG
TGCTACTCCTCCCTGCTGGACTCGATGAACGCTGAGATCCGCATCCGCTGGC
TGCAGATTGAGGTCCGCAACGACTACTATCCTGACCTCCACAGGGTGCGGCG
CTTCCTGGAGAGCCAGATGTCACGCATGTACACCATCCCGCTGTACGAGGAC
CTCTGCACCGGTGCCCTCAAGTCCTTCGCGCTGGAGGTCTTCTACCAGACGC
AGGGCCGGCTGCACCCCAACCTGCGCAGAGCCATCCAGCAGATCCTGTCCCA
GGGCCTGGGCTCCAGCACAGAGCCCGCCTCAGAGCCCAGCACGGAGCTGGGC
AAGGCTGAAGCAGACACAGACTCGGACGCACAGGCCCTGCTGCTTGGGGACG
AGGCCCCCAGCAGTGCCATCTCTCTCAGGACGTCAATGTGTCTGCCTAGCC
CTGTTGGCGGGCTGACCCTCGACCTCCCA GACACCACAATTGTGCCTTCTGT
GGGCCAGGCCTGCCATGACTGCGTCTCGGCTCTGGCCATGAGCTCTGCCCAG
GCCCACAAGCCCCTCCCCTGGGCTCTCCCAGGCAGGGAGAATGGGGAGAGGG
ACCTCCTTGTGTCTGGCAGAGACCTGTGGACCTGGCCTCCCCACTCCCAGCT
CTCTTGCACTGCAGGCCCTGGGGCCAGCCCGCACACACCATGCCTCCTGTCT
CAACACTGACAGCTGTGCCTAGCCCCGGATGCCAGCACCTGCCAGGTGCCGC
CCCGGGGCAAGGGCCCCAGCAGCCCTATGGTGACCGCCACACTGTGCCTTAA

TGTCTGCCGGGGGGCCCAGGCTGTGCTGTCCCTGCAGCACGCCTCCTTGCAAGG
GATCTGAGCCACCCTCCCCGCACAGCCCTGCACCCCGCCCCCTGGGGTTGGCA
GCCTCAGTTGGCCCCTGGCAGAGGAACAAGGACACAGACATTCCCTCAGTGT
GGGGGGCAGGGGACACAGGGAGAGGATGGTTGTCCCTGGGGAGGGCCCTCTG
GCCCCAGGCAACCTTAGCCCCCTCAGAACAGGGAGTCCCAGGACCCAGGGAGA
GTGTGGGGACAGGACAGCCTGTCTCTTGTAGCTTCCTGGGGTGGGAGGCACA
GGGGCAAAGCAATACCCCAGGGAAAGTGGGAGGTGGTGCTGGTGCTCTCTCC
AGGCCCACCATGCTGGGAGAGGCGGCCAGAGCCTGGGGCCTCCAGCCTGGGA
CTGCTGTGATGGGGTATCACGGTGATGGTCCCATTAACCTTCCACTCTGCAA
ACCTG

Fig. 2**SEQ ID NO: 2**

MWLSEGLATYAQRRITTETYGAAFTCLETAFRILDALHRQMKLLGEDSPVSKL
QVKLEPGVNP SHLMNLFTYEKGYCFVYYLSQLCGDPQRFDDFLRAYVEKYKF
TSVVAQDLLDSFLSFFPELKEQSVD CRAGLEFERWLNATGPPLAEPDLSQGS
SLTRPVEALFQLWTAEPLDQAAASAS AIDISKWRTFQTALFLDRLLDGSPLP
QEVVMSLSKCYSSLLDSMNAEIRIRWLQIEVRNDYYPDLHRVRRFLESQMSR
MYTIPLYEDLCTGALKSFALEV FYQTQGR LHPNLRRAIQQILSQGLGSSTEP
ASEPSTELGKAEADTDSDAQALLLGDEAPSSAISLRDVNVSA

Fig. 3**SEQ ID NO: 3**

5' -GGTCCGCAACGACTACTATCCT-3'

Fig. 4**SEQ ID NO: 4**

5' -TGGTGTACATGCGTGACATCTG-3'

Fig. 5

SEQ ID NO: 5

5' - CTCCACAGGGTGCGGCGCTTC - 3'